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<141> 2001-04-11

<150> PCT/US00/28664

<151> 2000-10-17

<150> 60/163,085

<151> 1999-11-02

<150> 60/172,411

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<170> PatentIn Ver. 2.0

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<213> Homo sapiens
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 <213> Homo sapiens

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692

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 <211> 700
 <212> DNA
 <213> Homo sapiens

<400> 26
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actttcaatt	gcctagcaca	atagtgaagta	catttgaatt	gaatatataa	taaatattgc	2700
aaaataaaaat	ccatctaaat	agaaaaaaa	aaaaaaaaa	aaaaaaactc	ga	2752

<210> 28

<211> 947

<212> DNA

<213> Homo sapiens

<400> 28

ggcagcagac	tggtactccc	ttgaatttat	aaacaaggag	gctcaaagag	ctgaagatta	60
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gccccaggtct	gactctaaaa	ttctgctttc	ttcatttagt	ttttctgcca	ttcaaagttt	180
actgccagct	cctccaggaa	cttctgagta	agcctgtctc	caaattgccc	ctaactcctc	240
agtgccagtc	atggggccagg	cccctggggg	acttggaatg	agtaaaacat	gcacctattg	300
ttaaaggttca	cctgggtccag	aagggggaggc	agaccacct	gccaccatg	acccggacag	360
aggcgataaaa	tacggagcac	ctcatggcca	gggcctggag	gattcccacc	ataccctacc	420
cagcgccaat	aacagagggg	ccctggagtt	gtgtacacaa	gatagcagtt	cagtgccttg	480
acacatacta	gttgagaagg	gaaatcaggg	ctaataaaac	agctcaactg	tctggcagag	540
ccaggctcct	tctgccatga	tacagtttgg	atgtttgtcc	ctttctaata	ttatatggaa	600
atttgatccc	cagtgttgaa	gggtggggcct	gggtgggagat	gttcctgtta	tcgggatgga	660
tccttcatga	atagcctggt	gccatctttg	tggttaacaag	tgagttctca	cttgattagt	720
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ctcttgctat	gtgatgtgc	tgccctttgc	cttccgtcat	gagtgggaagc	tcctgaagc	840
ccccaccaga	agcaaatgct	ggcaccatgc	tgctgttaca	gcctgcagaa	ctgtgagcca	900
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<210> 29

<211> 932

<212> DNA

<213> Homo sapiens

<400> 29

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aactctggat	ttgctgctgc	ctcttctgga	agccctgctg	gttttgggag	ttccccagca	180
tttgagactg	cagcctctac	cagttcaggt	atctctactt	ctgctccagc	ttttggattt	240
gggaagcctg	aagtcacatc	ggctgcatca	ttttcattca	aaagccctgc	agcttccagt	300
tttggatcac	ctggattttc	aggacttcca	gcttccttgg	caacagggtcc	gtcagagct	360
cccagtgggc	ccagcctttt	gaggtggcag	ttctgtggct	ggttttggta	gtccggggct	420
cacattctca	cactggcttt	ttctaagcca	tccagtggac	acttttggaa	atagccaggc	480
atatccctact	tctctgtcag	cccttcaagc	cagcatcaat	tgcaacagat	aatgtgttat	540

tcacacccag	aaataaacta	acagtagaag	aactggaaca	atttcaatcc	aagaaattta	600
ctctgggaaa	aattccatta	aagcctccac	ctctggaact	tctaaatgtt	taaaagggca	660
attttaata	caaaaaagaa	tgatgtttaa	aattgctttg	agtgattcat	acagagatgt	720
atatatgcat	acatgtatat	attcataagg	aatataagct	tccatcaata	gtgattttaa	780
atttgatatt	tttcttaact	ctaaatattt	aagtaaaaaag	taacaaaaaac	tctgcaagca	840
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<210> 30

<211> 1670

<212> DNA

<213> Homo sapiens

<400> 30

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aatggacggg	ttctgggaga	ccagatggct	tcagacactg	agctccagga	aatgtccacc	180
gaggggagta	agtacattaa	tcgggaaatt	aaaaatgctc	tcaagggggg	gaagcagata	240
aagacactaa	tagaacaac	aaacgaggag	cgcaaatccc	tgctcaccaa	cttgggaaga	300
gccaaaga	agaaagagga	tgccctgaat	gacaccaagg	attcagaaat	gaagctgaag	360
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gcaggacagt	ttcgaccggg	catccagcat	catggatgag	ctgttccagg	acagattctt	660
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tgcatgtcac	cgagtgcacg	ggccttcctt	gaggccctcc	tgtccctca	ccccgcctgt	1560
cctccctctg	gactctgcat	tgtaacaccg	tgttcactga	tcattgggaag	aactcctgtg	1620
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<210> 31

<211> 2072

<212> DNA

<213> Homo sapiens

<220>

<221> SITE

<222> (2065)

<223> n equals a,t,g, or c

<400> 31

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ggagcgcagg	gggcgcgcgg	cccggggact	cgcattcccc	ggttccccct	ccacccacg	120

cggcctggac	catggacgcc	agatggtggg	cagtgggtggt	gctggctgcg	ttccccctccc	180
taggggacagg	tggggagact	cccgaagccc	ctccggagtc	atggacccag	ctatgggttct	240
tccgatttgt	ggtgaatgct	gctggctatg	ccagctttat	ggtacctggc	tacctcctgg	300
tgcagtactt	caggcggaag	aactacctgg	agaccggtag	gggcctctgc	ttccccctgg	360
tgaaagcttg	tgtgtttggc	aatgagccca	aggcctctga	tgaggttccc	ctggcgcccc	420
gaacagaggc	ggcagagacc	accccgatgt	ggcaggccct	gaagctgctc	ttctgtgcca	480
cagggctcca	ggtgtcttat	ctgacttggg	gtgtgctgca	ggaaagagt	atgacccgca	540
gctatggggc	cacagccaca	tcaccgggtg	agcgctttac	ggactcgag	ttcctgggtg	600
taatgaaccg	agtgtctggc	ctgattgtgg	ctggcctctc	ctgtgttctc	tgcaagcagc	660
ccgggcatgg	ggcacccatg	taccgggtact	cctttgccag	cctgtccaat	gtgcttagca	720
gctggtgcca	atacgaagct	cttaagttcg	tcagcttccc	caccaggtg	ctggccaagg	780
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gttatattgc	ttttgacagc	ttcacctcaa	actggcagga	tgccctgttt	gcctataaga	1020
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gaagatgggt	ctgtgctgag	gaaaggggat	gcagagccct	gccagcacc	accacctcct	1920
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atgtaacttt	ttgtctttat	aattttattt	tattaaatta	aattactgca	aaaaaaaaaa	2040
aaaaaaamt	gggggggggg	ccggncccca	at			2072

<210> 32

<211> 985

<212> DNA

<213> Homo sapiens

<400> 32

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gcggaaaagg	acaaggatcc	aaactggcga	atttgctgat	cttcgcgtcc	ctctccgctt	180
tccggccggc	agcgtgcca	gggtatat	ccttttttcc	gacccctgca	cagcctcttt	240
aaactgttta	aatgagaatg	tccttggctc	agagagtact	actcacctgg	cttttcacac	300
tactcttctt	gatcatgttg	gtgttgaaac	tggatgagaa	agcaccttgg	aactggttcc	360
tcatattyat	tccagtctgg	atatttgata	ctatccttct	tgtcctgctg	attgtgaaaa	420
tggtggggcg	gtgtaagtct	ggctttgacc	ctcgacatgg	atcacacaat	attaaaaaaa	480
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gtacatcatc	tcctttctat	tgtgttcaa	caagttacca	ttaaagtgtt	ctgaatctgt	720
caagcttcaa	gaataccaga	gaactgaggg	aaaataccaa	atgtagtttt	atactacttc	780
cataaaacar	gattggtgaa	tcacggactt	ctagtcaacc	tacagcttaa	ttattcagca	840
tttgagttat	tgagatcctt	attatctcta	tgtaaataaa	gtttgttttg	gacctcaaaa	900
aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	960
accggggggg	ggggcccccc	cccaa				985

<210> 33
 <211> 1380
 <212> DNA
 <213> Homo sapiens

<400> 33
 tgaagacgcc ctccgtcagc gacgccgtcg caatggccat ttgtcaattc ttccttcaag 60
 gccggtgccg ctttggagat cgggtgctgga acgaacatcc cgggtgctagg ggtgcaggag 120
 gaggacggca gcaaccgcag cagcagcctt caggtaataa tagacgtgga tgggaatacaa 180
 ctagccagag atattccaat gtcattccagc catccagttt ctccaaatcc acaccatggg 240
 ggggcagcag agatcaagaa acctaataatt tcaggtttta cagacatttc accagaggaa 300
 ttgaggcttg aataccataa cttcttaacc agcaataact tacagagtta tctaaattct 360
 gtccaacgtt taataaatca atggaggaaac agggtaaatg aactgaaaag tctaaatata 420
 tcaactaaag tagctttgct ctctgatgta aaggatggag taaatccggc agcacctgca 480
 tttggatttg gcagcagtca agcagcaaca tttatgtcgc caggctttcc agtcaataac 540
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 tataagcttc catcaatagt gatttttaaat ttgatttttt tcttaactct aaatatattaa 1260
 gtaaaaagta acaaaaactc tgcaagcaag ggaatttttt tgtactgtaa ttttgaatgg 1320
 aactgaaaaa ttatgcacga ataaagtact tttctcatgc caaaaaaaaa aaaaaaaaaa 1380

<210> 34
 <211> 363
 <212> PRT
 <213> Homo sapiens

<400> 34
 Met Lys Thr Leu Leu Leu Leu Val Gly Leu Leu Leu Thr Trp Glu Asn
 1 5 10 15
 Gly Arg Val Leu Gly Asp Gln Met Val Ser Asp Thr Glu Leu Gln Glu
 20 25 30
 Met Ser Thr Glu Gly Ser Lys Tyr Ile Asn Arg Glu Ile Lys Asn Ala
 35 40 45
 Leu Lys Gly Val Lys Gln Ile Lys Thr Leu Ile Glu Gln Thr Asn Glu
 50 55 60
 Glu Arg Lys Ser Leu Leu Thr Asn Leu Glu Glu Ala Lys Lys Lys Lys
 65 70 75 80
 Glu Asp Ala Leu Asn Asp Thr Lys Asp Ser Glu Met Lys Leu Lys Ala
 85 90 95
 Ser Gln Gly Val Cys Asn Asp Thr Met Met Ala Leu Trp Glu Glu Cys

<210> 35

<211> 766

<212> PRT

<213> Homo sapiens

<400> 35

Met Ile Trp Arg Ser Arg Ala Gly Ala Glu Leu Phe Ser Leu Met Ala

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Leu Trp Glu Trp Ile Ala Leu Ser Leu His Cys Trp Val Leu Ala Val	20	25	30
Ala Ala Val Ser Asp Gln His Ala Thr Ser Pro Phe Asp Trp Leu Leu	35	40	45
Ser Asp Lys Gly Pro Phe His Arg Ser Gln Glu Tyr Thr Asp Phe Val	50	55	60
Asp Arg Ser Arg Gln Gly Phe Ser Thr Arg Tyr Lys Ile Tyr Arg Glu	65	70	75
Phe Gly Arg Trp Lys Val Asn Asn Leu Ala Val Glu Arg Arg Asn Phe	85	90	95
Leu Gly Ser Pro Leu Pro Leu Ala Pro Glu Phe Phe Arg Asn Ile Arg	100	105	110
Leu Leu Gly Arg Arg Pro Thr Leu Gln Gln Ile Thr Glu Asn Leu Ile	115	120	125
Lys Lys Tyr Gly Thr His Phe Leu Leu Ser Ala Thr Leu Gly Gly Glu	130	135	140
Glu Ser Leu Thr Ile Phe Val Asp Lys Arg Lys Leu Ser Lys Arg Ala	145	150	155
Glu Gly Ser Asp Ser Thr Thr Asn Ser Ser Ser Val Thr Leu Glu Thr	165	170	175
Leu His Gln Leu Ala Ala Ser Tyr Phe Ile Asp Arg Asp Ser Thr Leu	180	185	190
Arg Arg Leu His His Ile Gln Ile Ala Ser Thr Ala Ile Lys Val Thr	195	200	205
Glu Thr Arg Thr Gly Pro Leu Gly Cys Ser Asn Tyr Asp Asn Leu Asp	210	215	220
Ser Val Ser Ser Val Leu Val Gln Ser Pro Glu Asn Lys Ile Gln Leu	225	230	235
Gln Gly Leu Gln Val Leu Leu Pro Asp Tyr Leu Gln Glu Arg Phe Val	245	250	255
Gln Ala Ala Leu Ser Tyr Ile Ala Cys Asn Ser Glu Gly Glu Phe Ile	260	265	270
Cys Lys Glu Asn Asp Cys Trp Cys His Cys Gly Pro Lys Phe Pro Glu	275	280	285
Cys Asn Cys Pro Ser Met Asp Ile Gln Ala Met Glu Glu Asn Leu Leu	290	295	300
Arg Ile Thr Glu Thr Trp Lys Ala Tyr Asn Ser Asp Phe Glu Glu Ser	305	310	315
			320

Asp	Glu	Phe	Lys	Leu	Phe	Met	Lys	Arg	Leu	Pro	Met	Asn	Tyr	Phe	Leu	
				325					330						335	
Asn	Thr	Ser	Thr	Ile	Met	His	Leu	Trp	Thr	Met	Asp	Ser	Asn	Phe	Gln	
			340					345					350			
Arg	Arg	Tyr	Glu	Gln	Leu	Glu	Asn	Ser	Met	Lys	Gln	Leu	Phe	Leu	Lys	
		355					360					365				
Ala	Gln	Lys	Ile	Val	His	Lys	Leu	Phe	Ser	Leu	Ser	Lys	Arg	Cys	His	
	370					375					380					
Lys	Gln	Pro	Leu	Ile	Ser	Leu	Pro	Arg	Gln	Arg	Thr	Ser	Thr	Tyr	Trp	
385					390					395					400	
Leu	Thr	Arg	Ile	Gln	Ser	Phe	Leu	Tyr	Cys	Asn	Glu	Asn	Gly	Leu	Leu	
			405						410					415		
Gly	Ser	Phe	Ser	Glu	Glu	Thr	His	Ser	Cys	Thr	Cys	Pro	Asn	Asp	Gln	
			420					425					430			
Val	Val	Cys	Thr	Ala	Phe	Leu	Pro	Cys	Thr	Val	Gly	Asp	Ala	Ser	Ala	
		435					440					445				
Cys	Leu	Thr	Cys	Ala	Pro	Asp	Asn	Arg	Thr	Arg	Cys	Gly	Thr	Cys	Asn	
	450					455					460					
Thr	Gly	Tyr	Met	Leu	Ser	Gln	Gly	Leu	Cys	Lys	Pro	Glu	Val	Ala	Glu	
465					470					475					480	
Ser	Thr	Asp	His	Tyr	Ile	Gly	Phe	Glu	Thr	Asp	Leu	Gln	Asp	Leu	Glu	
			485						490					495		
Met	Lys	Tyr	Leu	Leu	Gln	Lys	Thr	Asp	Arg	Arg	Ile	Glu	Val	His	Ala	
			500					505					510			
Ile	Phe	Ile	Ser	Asn	Asp	Met	Arg	Leu	Asn	Ser	Trp	Phe	Asp	Pro	Ser	
		515					520					525				
Trp	Arg	Lys	Arg	Met	Leu	Leu	Thr	Leu	Lys	Ser	Asn	Lys	Tyr	Lys	Ser	
	530					535					540					
Ser	Leu	Val	His	Met	Ile	Leu	Gly	Leu	Ser	Leu	Gln	Ile	Cys	Leu	Thr	
545					550					555					560	
Lys	Asn	Ser	Thr	Leu	Glu	Pro	Val	Leu	Ala	Val	Tyr	Val	Asn	Pro	Phe	
				565					570					575		
Gly	Gly	Ser	His	Ser	Glu	Ser	Trp	Phe	Met	Pro	Val	Asn	Glu	Asn	Ser	
			580					585					590			
Phe	Pro	Asp	Trp	Glu	Arg	Thr	Lys	Leu	Asp	Leu	Pro	Leu	Gln	Cys	Tyr	
		595					600					605				
Asn	Trp	Thr	Leu	Thr	Leu	Gly	Asn	Lys	Trp	Lys	Thr	Phe	Phe	Glu	Thr	
	610					615					620					

Lys Ser Thr Pro Lys Thr Thr Ser Val Ser Gln Asn Thr Ser Gln Ile
115 120 125

Ser Thr Ser Thr Met Thr Val Thr His Asn Ser Ser Val Thr Ser Ala
130 135 140

Ala Ser Ser Val Thr Ile Thr Thr Thr Met His Ser Glu Ala Lys Lys
145 150 155 160

Gly Ser Lys Phe Asp Thr Gly Ser Phe Val Gly Gly Ile Val Leu Thr
165 170 175

Leu Gly Val Leu Ser Ile Leu Tyr Ile Gly Cys Lys Met Tyr Tyr Ser
180 185 190

Arg Arg Gly Ile Arg Tyr Arg Thr Ile Asp Glu His Asp Ala Ile Ile
195 200 205

<210> 37
<211> 605
<212> PRT
<213> Homo sapiens

<400> 37

Met Gly Arg Leu Leu Arg Ala Ala Arg Leu Pro Pro Leu Leu Ser Pro
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Leu Leu Leu Leu Leu Val Gly Gly Ala Phe Leu Gly Ala Cys Val Ala
20 25 30

Gly Ser Asp Glu Pro Gly Pro Glu Gly Leu Thr Ser Thr Ser Leu Leu
35 40 45

Asp Leu Leu Leu Pro Thr Gly Leu Glu Pro Leu Asp Ser Glu Glu Pro
50 55 60

Ser Glu Thr Met Gly Leu Gly Ala Gly Leu Gly Ala Pro Gly Ser Gly
65 70 75 80

Phe Pro Ser Glu Glu Asn Glu Glu Ser Arg Ile Leu Gln Pro Pro Gln
85 90 95

Tyr Phe Trp Glu Glu Glu Glu Glu Leu Asn Asp Ser Ser Leu Asp Leu
100 105 110

Gly Pro Thr Ala Asp Tyr Val Phe Pro Asp Leu Thr Glu Lys Ala Gly
115 120 125

Ser Ile Glu Asp Thr Ser Gln Ala Gln Glu Leu Pro Asn Leu Pro Ser
130 135 140

Pro Leu Pro Lys Met Asn Leu Val Glu Pro Pro Trp His Met Pro Pro
145 150 155 160

Arg Glu Glu Glu Glu Glu Glu Glu Glu Glu Glu Arg Glu Lys Glu
165 170 175

04110-02120000

Glu	Val	Glu	Lys	Gln	Glu	Glu	Glu	Glu	Glu	Glu	Glu	Leu	Leu	Pro	Val
			180					185					190		
Asn	Gly	Ser	Gln	Glu	Glu	Ala	Lys	Pro	Gln	Val	Arg	Asp	Phe	Ser	Leu
		195					200					205			
Thr	Ser	Ser	Ser	Gln	Thr	Pro	Gly	Ala	Thr	Lys	Ser	Arg	His	Glu	Asp
	210					215					220				
Ser	Gly	Asp	Gln	Ala	Ser	Ser	Gly	Val	Glu	Val	Glu	Ser	Ser	Met	Gly
225					230					235					240
Pro	Ser	Leu	Leu	Leu	Pro	Ser	Val	Thr	Pro	Thr	Thr	Val	Thr	Pro	Gly
				245					250					255	
Asp	Gln	Asp	Ser	Thr	Ser	Gln	Glu	Ala	Glu	Ala	Thr	Val	Leu	Pro	Ala
			260					265					270		
Ala	Gly	Leu	Gly	Val	Glu	Phe	Glu	Ala	Pro	Gln	Glu	Ala	Ser	Glu	Glu
		275					280					285			
Ala	Thr	Ala	Gly	Ala	Ala	Gly	Leu	Ser	Gly	Gln	His	Glu	Glu	Val	Pro
	290					295					300				
Ala	Leu	Pro	Ser	Phe	Pro	Gln	Thr	Thr	Ala	Pro	Ser	Gly	Ala	Glu	His
305					310					315					320
Pro	Asp	Glu	Asp	Pro	Leu	Gly	Ser	Arg	Thr	Ser	Ala	Ser	Ser	Pro	Leu
				325					330					335	
Ala	Pro	Gly	Asp	Met	Glu	Leu	Thr	Pro	Ser	Ser	Ala	Thr	Leu	Gly	Gln
			340					345					350		
Glu	Asp	Leu	Asn	Gln	Gln	Leu	Leu	Glu	Gly	Gln	Ala	Ala	Glu	Ala	Gln
		355					360					365			
Ser	Arg	Ile	Pro	Trp	Asp	Ser	Thr	Gln	Val	Ile	Cys	Lys	Asp	Trp	Ser
	370				375						380				
Asn	Leu	Ala	Gly	Lys	Asn	Tyr	Ile	Ile	Leu	Asn	Met	Thr	Glu	Asn	Ile
385					390					395					400
Asp	Cys	Glu	Val	Phe	Arg	Gln	His	Arg	Gly	Pro	Gln	Leu	Leu	Ala	Leu
				405					410					415	
Val	Glu	Glu	Val	Leu	Pro	Arg	His	Gly	Ser	Gly	His	His	Gly	Ala	Trp
			420					425					430		
His	Ile	Ser	Leu	Ser	Lys	Pro	Ser	Glu	Lys	Glu	Gln	His	Leu	Leu	Met
		435					440					445			
Thr	Leu	Val	Gly	Glu	Gln	Gly	Val	Val	Pro	Thr	Gln	Asp	Val	Leu	Ser
		450				455					460				
Met	Leu	Gly	Asp	Ile	Arg	Arg	Ser	Leu	Glu	Glu	Ile	Gly	Ile	Gln	Asn
465					470					475					480
Tyr	Ser	Thr	Thr	Ser	Ser	Cys	Gln	Ala	Arg	Ala	Ser	Gln	Val	Arg	Ser

495

Met Ala Ala Ala Ser Ala Gly Ala Thr Arg Leu Leu Leu Leu Leu Leu
1 5 10 15

Leu Gly Ala Gly Gly Glu Thr Pro Glu Ala Pro Pro Glu Ser Trp Thr
20 25 30

Gln	Leu	Trp	Phe	Phe	Arg	Phe	Val	Val	Asn	Ala	Ala	Gly	Tyr	Ala	Ser
		35					40					45			
Phe	Met	Val	Pro	Gly	Tyr	Leu	Leu	Val	Gln	Tyr	Phe	Arg	Arg	Lys	Asn
	50					55					60				
Tyr	Leu	Glu	Thr	Gly	Arg	Gly	Leu	Cys	Phe	Pro	Leu	Val	Lys	Ala	Cys
65					70					75					80
Val	Phe	Gly	Asn	Glu	Pro	Lys	Ala	Ser	Asp	Glu	Val	Pro	Leu	Ala	Pro
				85					90					95	
Arg	Thr	Glu	Ala	Ala	Glu	Thr	Thr	Pro	Met	Trp	Gln	Ala	Leu	Lys	Leu
			100					105					110		
Leu	Phe	Cys	Ala	Thr	Gly	Leu	Gln	Val	Ser	Tyr	Leu	Thr	Trp	Gly	Val
		115					120					125			
Leu	Gln	Glu	Arg	Val	Met	Thr	Arg	Ser	Tyr	Gly	Ala	Thr	Ala	Thr	Ser
		130				135					140				
Pro	Gly	Glu	Arg	Phe	Thr	Asp	Ser	Gln	Phe	Leu	Val	Leu	Met	Asn	Arg
145					150					155					160
Val	Leu	Ala	Leu	Ile	Val	Ala	Gly	Leu	Ser	Cys	Val	Leu	Cys	Lys	Gln
				165				170						175	
Pro	Arg	His	Gly	Ala	Pro	Met	Tyr	Arg	Tyr	Ser	Phe	Ala	Ser	Leu	Ser
			180					185					190		
Asn	Val	Leu	Ser	Ser	Trp	Cys	Gln	Tyr	Glu	Ala	Leu	Lys	Phe	Val	Ser
		195					200					205			
Phe	Pro	Thr	Gln	Val	Leu	Ala	Lys	Ala	Ser	Lys	Val	Ile	Pro	Val	Met
		210				215					220				
Leu	Met	Gly	Lys	Leu	Val	Ser	Arg	Arg	Ser	Tyr	Glu	His	Trp	Glu	Tyr
225					230					235					240
Leu	Thr	Ala	Thr	Leu	Ile	Ser	Ile	Gly	Val	Ser	Met	Phe	Leu	Leu	Ser
				245					250					255	
Ser	Gly	Pro	Glu	Pro	Arg	Ser	Ser	Pro	Ala	Thr	Thr	Leu	Ser	Gly	Leu
			260					265					270		
Ile	Leu	Leu	Ala	Gly	Tyr	Ile	Ala	Phe	Asp	Ser	Phe	Thr	Ser	Asn	Trp
		275					280					285			
Gln	Asp	Ala	Leu	Phe	Ala	Tyr	Lys	Met	Ser	Ser	Val	Gln	Met	Met	Phe
		290				295					300				
Gly	Val	Asn	Phe	Phe	Ser	Cys	Leu	Phe	Thr	Val	Gly	Ser	Leu	Leu	Glu
305					310					315					320
Gln	Gly	Ala	Leu	Leu	Glu	Gly	Thr	Arg	Phe	Met	Gly	Arg	His	Ser	Glu
				325				330						335	

<213> Homo sapiens

Met Arg Leu Pro Ala Trp Cys Arg His Thr Thr Leu Ala Ile Ser Cys
1 5 10 15

Pro Thr Ile Ser His Leu Gly Val Lys Pro Leu Ser Val Gly Trp Gly
35 40 45

Ala Pro Ser Thr Leu Pro Val Ser Pro Cys Gly Gly Lys Pro Ala Ala
50 55 60

Pro Thr Ser Ala Ser Pro Ala Ala Ala Pro Leu Arg Phe Trp Arg Pro
65 70 75 80

Gly Ala Ser Gly Gly Gly Ala Gly Gly Thr Arg Arg Leu Ala Leu Cys
85 90 95

Arg Leu Val Thr Ala Arg Thr Thr Leu Ala Thr Gly Thr Pro Gly Leu
100 105 110

Ser Ala Arg Pro Arg Gln Arg Pro Cys Leu Leu Pro Val Leu Pro Arg
115 120 125

Arg Pro Ala Glu Leu Ser Val Ser Leu Glu Pro Ser Pro Gly Ser Ser
130 135 140

Gly Arg Gly Phe Leu Cys Leu Pro Phe Cys Lys Arg Asp Ala Asp Thr
145 150 155 160

Ser Leu Gly Gln Thr Leu Thr Ser Ser Cys Ser Leu Ser Ser Ile Leu
165 170 175

Val Gly Gly Thr Leu Arg Pro Arg Cys Ser Cys Pro Pro Phe Thr Gln
180 185 190

Arg Ser Ala Phe His Leu Arg Thr Pro His Asn Gln Tyr His His Gly
195 200 205

Ser Thr Ser Leu Ala Ser His
210 215

<213> Homo sapiens

Met Lys Ser Ala Leu His Arg Asp Ile Cys Ile Leu Met Leu Thr Ala
1 5 10 15

Ala Leu Phe Thr Ile Ala Lys Thr Glu Lys Gln His Lys Cys Pro Ser
20 25 30

Ile Asp Glu Gln Ile Asn Asn Leu Gln Tyr Ile Cys Thr Met Glu Tyr
 35 40 45

His Ser Ala Leu Gln Lys Glu Met Leu Leu Tyr Leu Gln
 50 55 60

<210> 45
 <211> 125
 <212> PRT
 <213> Homo sapiens

<400> 45
 Met Ile Pro Phe Pro Ala Cys Leu Leu Leu Ala Leu Phe Pro Lys Val
 1 5 10 15

Gln Val Gly Arg Thr Thr Ser Ala Tyr Phe Ser Thr Ile Pro Ser Met
 20 25 30

Pro Ala Arg Ser Gln Ile Asn Leu Pro Val Glu Ser Gly Ser Ala Leu
 35 40 45

Leu Glu Pro Arg Gly Lys Gly Arg Val Glu Arg Val Cys Pro Val Ala
 50 55 60

Trp Ser Ser Met Val Ala Ser Cys Leu Pro Ser Pro Ser Ser Gly Gly
 65 70 75 80

Pro Glu Gly Ser Leu Gly Thr Val Pro Gln Ile Leu Thr Gln Gly Pro
 85 90 95

Ala Trp Gly Arg Asp Gly Cys Arg Gln Asn Ala Leu Tyr Arg Asp Phe
 100 105 110

Leu Leu Leu Gly Arg Cys Val Ser Pro Thr Ile Cys Leu
 115 120 125

<210> 46
 <211> 71
 <212> PRT
 <213> Homo sapiens

<400> 46
 Met Leu Val Ala Ala Ile Val Phe Ile Ser Phe Gly Val Val Ala Ala
 1 5 10 15

Phe Cys Cys Ala Ile Val Asp Gly Val Phe Ala Ala Gln His Ile Glu
 20 25 30

Pro Lys Ala Pro His His Gly Lys Met Pro Val Tyr Ser Ser Gly Val
 35 40 45

Gly Tyr Leu Tyr Asp Val Tyr Gln Thr Glu Val Ser Arg Ser Thr Glu
 50 55 60

Ile His Val Gly Leu Leu Asn

70

<211> 69

<212> PRT

<213> Homo sapiens

<400> 47

Met Lys Ala Val Val Leu Leu Lys Ala Phe Ser Phe Ser Leu Cys Ser
1 5 10 15

Ala Ile Ser Pro Val Thr Pro Gly Phe Arg Gln Thr Ile Asn Val Leu
20 25 30

Asp Thr Val Ala Phe Ser Ala Phe Phe Ile Tyr Leu Phe Thr Val Thr
35 40 45

Ala Ser Ile Asn Phe Tyr Ala Tyr Phe Ser Ser Phe Leu Ala Gly Ala
50 55 60

Pro Phe Ile Lys Ile
65

<210> 48

<211> 85

<212> PRT

<213> Homo sapiens

<400> 48

Met Ala Ala Gly Gly Cys Leu Leu Leu Leu Ala Phe Phe Pro Leu Ser
1 5 10 15

Arg Gly Ser His Phe His Leu Gln Lys Arg Ala Leu Ala Glu Ala Ser
20 25 30

Phe Glu Ala Thr Leu Cys Glu Leu Phe Val Ile Glu Thr Ala Ser Lys
35 40 45

Gly Thr Leu Leu Ile Ile Thr Ile Arg His Leu Val Thr Tyr Ile Ile
50 55 60

Val Ile Phe Lys Cys His Met Leu Lys Asn Glu Met Asn Ser Ser Ile
65 70 75 80

Lys Pro His Phe Gln
85

<210> 49

<211> 150

<212> PRT

<213> Homo sapiens

<400> 49

Met Val Met Ile Leu Phe Val Ala Phe Ile Thr Cys Trp Glu Glu Val
1 5 10 15

Thr	Thr	Leu	Val	Gln	Ala	Ile	Arg	Ile	Thr	Ser	Tyr	Met	Asn	Glu	Thr
			20					25					30		
Ile	Leu	Tyr	Phe	Pro	Phe	Ser	Ser	His	Ser	Ser	Tyr	Thr	Val	Arg	Ser
		35					40					45			
Lys	Lys	Ile	Phe	Leu	Ser	Lys	Leu	Ile	Val	Cys	Phe	Leu	Ser	Thr	Trp
		50				55					60				
Leu	Pro	Phe	Val	Leu	Leu	Gln	Val	Ile	Ile	Val	Leu	Leu	Lys	Val	Gln
65					70					75					80
Ile	Pro	Ala	Tyr	Ile	Glu	Met	Asn	Ile	Pro	Trp	Leu	Tyr	Phe	Val	Asn
				85					90					95	
Ser	Phe	Leu	Ile	Ala	Thr	Val	Tyr	Trp	Phe	Asn	Cys	His	Lys	Leu	Asn
			100					105					110		
Leu	Lys	Asp	Ile	Gly	Leu	Pro	Leu	Asp	Pro	Phe	Val	Asn	Trp	Lys	Cys
		115					120					125			
Cys	Phe	Ile	Pro	Leu	Thr	Ile	Pro	Asn	Leu	Glu	Gln	Ile	Glu	Lys	Pro
		130				135					140				
Ile	Ser	Ile	Met	Ile	Cys										
145					150										
<210> 50															
<211> 298															
<212> PRT															
<213> Homo sapiens															
<400> 50															
Met	Lys	Thr	Leu	Gln	Ser	Thr	Leu	Leu	Leu	Leu	Leu	Leu	Val	Pro	Leu
1				5					10					15	
Ile	Lys	Pro	Ala	Pro	Pro	Thr	Gln	Gln	Asp	Ser	Arg	Ile	Ile	Tyr	Asp
			20					25					30		
Tyr	Gly	Thr	Asp	Asn	Phe	Glu	Glu	Ser	Ile	Phe	Ser	Gln	Asp	Tyr	Glu
		35					40					45			
Asp	Lys	Tyr	Leu	Asp	Gly	Lys	Asn	Ile	Lys	Glu	Lys	Glu	Thr	Val	Ile
	50					55					60				
Ile	Pro	Asn	Glu	Lys	Ser	Leu	Gln	Leu	Gln	Lys	Asp	Glu	Ala	Ile	Thr
65					70					75					80
Pro	Leu	Pro	Pro	Lys	Lys	Glu	Asn	Asp	Glu	Met	Pro	Thr	Cys	Leu	Leu
				85					90					95	
Cys	Val	Cys	Leu	Ser	Gly	Ser	Val	Tyr	Cys	Glu	Glu	Val	Asp	Ile	Asp
			100					105					110		
Ala	Val	Pro	Pro	Leu	Pro	Lys	Glu	Ser	Ala	Tyr	Leu	Tyr	Ala	Arg	Phe
			115				120					125			

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<210> 52
<211> 145
<212> PRT
<213> Homo sapiens
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Met Leu Arg Thr Leu Val Leu Lys Gln Thr Leu Asp Leu Leu Leu Pro
1 5 10 15

Leu Leu Glu Ala Leu Leu Val Leu Gly Val Pro Gln His Leu Glu Leu
20 25 30

Gln Pro Leu Pro Val Gln Val Ser Leu Leu Leu Leu Gln Leu Leu Asp
35 40 45

Leu Gly Ser Leu Lys Ser His Arg Leu His His Phe His Ser Lys Ala
50 55 60

Leu Gln Leu Pro Val Leu Asp His Leu Asp Phe Gln Asp Phe Gln Leu
65 70 75 80

Pro Trp Gln Gln Val Leu Ser Glu Leu Pro Val Ala Pro Ala Phe Gly
85 90 95

Gly Gly Ser Ser Val Ala Gly Phe Gly Ser Pro Gly Leu Thr Phe Ser
100 105 110

His Trp Leu Phe Leu Ser His Pro Val Asp Thr Phe Gly Asn Ser Gln
115 120 125

Ala Tyr Pro Thr Ser Leu Ser Ala Leu Gln Ala Ser Ile Asn Cys Asn
130 135 140

Arg
145

<210> 53

<211> 139

<212> PRT

<213> Homo sapiens

<400> 53

Met Lys Thr Leu Leu Leu Leu Val Gly Leu Leu Leu Thr Trp Glu Asn
1 5 10 15

Gly Arg Val Leu Gly Asp Gln Met Val Ser Asp Thr Glu Leu Gln Glu
20 25 30

Met Ser Thr Glu Gly Ser Lys Tyr Ile Asn Arg Glu Ile Lys Asn Ala
35 40 45

Leu Lys Gly Val Lys Gln Ile Lys Thr Leu Ile Glu Gln Thr Asn Glu
50 55 60

Glu Arg Lys Ser Leu Leu Thr Asn Leu Glu Glu Ala Lys Lys Lys Lys
65 70 75 80

Glu Asp Ala Leu Asn Asp Thr Lys Asp Ser Glu Met Lys Leu Lys Ala
85 90 95

Ser Gln Gly Val Cys Asn Asp Thr Met Met Ala Leu Trp Glu Glu Cys

100 105 110
 Lys Pro Cys Leu Lys Gln Thr Trp Gly Lys Gly Leu Arg Pro Ser Leu
 115 120 125
 Gln Lys Gln His Arg Ala Gly Trp Pro Pro Gly
 130 135

 <210> 54
 <211> 432
 <212> PRT
 <213> Homo sapiens

 <400> 54
 Met Asp Ala Arg Trp Trp Ala Val Val Val Leu Ala Ala Phe Pro Ser
 1 5 10 15
 Leu Gly Ala Gly Gly Glu Thr Pro Glu Ala Pro Pro Glu Ser Trp Thr
 20 25 30
 Gln Leu Trp Phe Phe Arg Phe Val Val Asn Ala Ala Gly Tyr Ala Ser
 35 40 45
 Phe Met Val Pro Gly Tyr Leu Leu Val Gln Tyr Phe Arg Arg Lys Asn
 50 55 60
 Tyr Leu Glu Thr Gly Arg Gly Leu Cys Phe Pro Leu Val Lys Ala Cys
 65 70 75 80
 Val Phe Gly Asn Glu Pro Lys Ala Ser Asp Glu Val Pro Leu Ala Pro
 85 90 95
 Arg Thr Glu Ala Ala Glu Thr Thr Pro Met Trp Gln Ala Leu Lys Leu
 100 105 110
 Leu Phe Cys Ala Thr Gly Leu Gln Val Ser Tyr Leu Thr Trp Gly Val
 115 120 125
 Leu Gln Glu Arg Val Met Thr Arg Ser Tyr Gly Ala Thr Ala Thr Ser
 130 135 140
 Pro Gly Glu Arg Phe Thr Asp Ser Gln Phe Leu Val Leu Met Asn Arg
 145 150 155 160
 Val Leu Ala Leu Ile Val Ala Gly Leu Ser Cys Val Leu Cys Lys Gln
 165 170 175
 Pro Arg His Gly Ala Pro Met Tyr Arg Tyr Ser Phe Ala Ser Leu Ser
 180 185 190
 Asn Val Leu Ser Ser Trp Cys Gln Tyr Glu Ala Leu Lys Phe Val Ser
 195 200 205
 Phe Pro Thr Gln Val Leu Ala Lys Ala Ser Lys Val Ile Pro Val Met
 210 215 220
 Leu Met Gly Lys Leu Val Ser Arg Arg Ser Tyr Glu His Trp Glu Tyr

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225 230 235 240
 Leu Thr Ala Thr Leu Ile Ser Ile Gly Val Ser Met Phe Leu Leu Ser
 245 250 255
 Ser Gly Pro Glu Pro Arg Ser Ser Pro Ala Thr Thr Leu Ser Gly Leu
 260 265 270
 Ile Leu Leu Ala Gly Tyr Ile Ala Phe Asp Ser Phe Thr Ser Asn Trp
 275 280 285
 Gln Asp Ala Leu Phe Ala Tyr Lys Met Ser Ser Val Gln Met Met Phe
 290 295 300
 Gly Val Asn Phe Phe Ser Cys Leu Phe Thr Val Gly Ser Leu Leu Glu
 305 310 315 320
 Gln Gly Ala Leu Leu Glu Gly Thr Arg Phe Met Gly Arg His Ser Glu
 325 330 335
 Phe Ala Ala His Ala Leu Leu Leu Ser Ile Cys Ser Ala Cys Gly Gln
 340 345 350
 Leu Phe Ile Phe Tyr Thr Ile Gly Gln Phe Gly Ala Ala Val Phe Thr
 355 360 365
 Ile Ile Met Thr Leu Arg Gln Ala Phe Ala Ile Leu Leu Ser Cys Leu
 370 375 380
 Leu Tyr Gly His Thr Val Thr Val Val Gly Gly Leu Gly Val Ala Val
 385 390 395 400
 Val Phe Ala Ala Leu Leu Leu Arg Val Tyr Ala Arg Gly Arg Leu Lys
 405 410 415
 Gln Arg Gly Lys Lys Ala Val Pro Val Glu Ser Pro Val Gln Lys Val
 420 425 430

<210> 55

<211> 133

<212> PRT

<213> Homo sapiens

<400> 55

Met Arg Met Ser Leu Ala Gln Arg Val Leu Leu Thr Trp Leu Phe Thr
 1 5 10 15
 Leu Leu Phe Leu Ile Met Leu Val Leu Lys Leu Asp Glu Lys Ala Pro
 20 25 30
 Trp Asn Trp Phe Leu Ile Phe Ile Pro Val Trp Ile Phe Asp Thr Ile
 35 40 45
 Leu Leu Val Leu Leu Ile Val Lys Met Ala Gly Arg Cys Lys Ser Gly

50 55 60
 Phe Asp Pro Arg His Gly Ser His Asn Ile Lys Lys Lys Ala Trp Tyr
 65 70 75 80
 Leu Ile Ala Met Leu Leu Lys Leu Ala Phe Cys Leu Ala Leu Cys Ala
 85 90 95
 Lys Leu Glu Gln Phe Thr Thr Met Asn Leu Ser Tyr Val Phe Ile Pro
 100 105 110
 Leu Trp Ala Leu Leu Ala Gly Ala Leu Thr Glu Leu Gly Tyr Asn Val
 115 120 125
 Phe Phe Val Arg Asp
 130

<210> 56
 <211> 77
 <212> PRT
 <213> Homo sapiens

<400> 56
 Met Ala Ile Cys Gln Phe Phe Leu Gln Gly Arg Cys Arg Phe Gly Asp
 1 5 10 15
 Arg Cys Trp Asn Glu His Pro Gly Ala Arg Gly Ala Gly Gly Gly Arg
 20 25 30
 Gln Gln Pro Gln Gln Gln Pro Ser Gly Asn Asn Arg Arg Gly Trp Asn
 35 40 45
 Thr Thr Ser Gln Arg Tyr Ser Asn Val Ile Gln Pro Ser Ser Phe Ser
 50 55 60
 Lys Ser Thr Pro Trp Gly Gly Ser Arg Asp Gln Glu Thr
 65 70 75

<210> 57
 <211> 247
 <212> PRT
 <213> Homo sapiens

<400> 57
 Asn Arg Pro Gly Gly Arg Val Tyr Ala Arg Val Cys Arg Ser Ser Thr
 1 5 10 15
 Gly Leu Val Gly His Gln Val Glu Glu Phe Leu Asn Gln Ser Ser Pro
 20 25 30
 Phe Tyr Phe Trp Ile Asn Gly Asp Arg Ile Asp Ser Leu Leu Glu Asn
 35 40 45
 Asp Arg Gln Gln Thr His Ala Leu Asp Val Met Gln Asp Ser Phe Asp
 50 55 60

Thr Lys Ile Gly Arg Thr Arg Ala Val Gly Lys Met Ser Ser Ser Leu
65 70 75 80

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<210> 59
<211> 468
<212> PRT
<213> Homo sapiens
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<400> 59															
His 1	Glu	Gly	Ser	Leu 5	Ala	Ala	Pro	Gly	Gly 10	Gly	Gly	Gly	Ser	Ala 15	Gly
Gly	Ala	Arg	Pro 20	Gly	Asp	Ser	His	Ser 25	Pro	Val	Pro	Pro	Pro 30	Pro	His
Ala	Ala	Trp 35	Thr	Met	Asp	Ala	Arg 40	Trp	Trp	Ala	Val	Val 45	Val	Leu	Ala
Ala	Phe 50	Pro	Ser	Leu	Gly	Ala 55	Gly	Gly	Glu	Thr	Pro 60	Glu	Ala	Pro	Pro
Glu 65	Ser	Trp	Thr	Gln	Leu 70	Trp	Phe	Phe	Arg	Phe 75	Val	Val	Asn	Ala	Ala 80
Gly	Tyr	Ala	Ser	Phe 85	Met	Val	Pro	Gly	Tyr 90	Leu	Leu	Val	Gln	Tyr 95	Phe
Arg	Arg	Lys	Asn 100	Tyr	Leu	Glu	Thr	Gly 105	Arg	Gly	Leu	Cys	Phe 110	Pro	Leu
Val	Lys	Ala 115	Cys	Val	Phe	Gly	Asn 120	Glu	Pro	Lys	Ala	Ser 125	Asp	Glu	Val
Pro	Leu 130	Ala	Pro	Arg	Thr	Glu 135	Ala	Ala	Glu	Thr	Thr 140	Pro	Met	Trp	Gln
Ala 145	Leu	Lys	Leu	Leu	Phe 150	Cys	Ala	Thr	Gly	Leu 155	Gln	Val	Ser	Tyr	Leu 160
Thr	Trp	Gly	Val	Leu 165	Gln	Glu	Arg	Val	Met 170	Thr	Arg	Ser	Tyr	Gly 175	Ala
Thr	Ala	Thr	Ser 180	Pro	Gly	Glu	Arg	Phe 185	Thr	Asp	Ser	Gln	Phe 190	Leu	Val
Leu	Met	Asn 195	Arg	Val	Leu	Ala	Leu 200	Ile	Val	Ala	Gly	Leu 205	Ser	Cys	Val
Leu 210	Cys	Lys	Gln	Pro	Arg	His 215	Gly	Ala	Pro	Met	Tyr 220	Arg	Tyr	Ser	Phe
Ala 225	Ser	Leu	Ser	Asn	Val 230	Leu	Ser	Ser	Trp	Cys 235	Gln	Tyr	Glu	Ala	Leu 240
Lys	Phe	Val	Ser	Phe 245	Pro	Thr	Gln	Val	Leu 250	Ala	Lys	Ala	Ser	Lys 255	Val

Trp Asn Trp Phe Leu Ile Phe Ile Pro Val Trp Ile Phe Asp Thr Ile
35 40 45

Leu Leu Val Leu Leu Ile Val Lys Met Ala Gly Arg Cys Lys Ser Gly
 50 55 60
 Phe Asp Pro Arg His Gly Ser His Asn Ile Lys Lys Lys Ala Trp Tyr
 65 70 75 80
 Leu Ile Ala Met Leu Leu Lys Leu Ala Phe Cys Leu Ala Leu Cys Ala
 85 90 95
 Lys Leu Glu Gln Phe Thr Thr Met Asn Leu Ser Tyr Val Phe Ile Pro
 100 105 110
 Leu Trp Ala Leu Leu Ala Gly Ala Leu Thr Glu Leu Gly Tyr Asn Val
 115 120 125
 Phe Phe Val Arg Asp
 130

<210> 61
 <211> 75
 <212> PRT
 <213> Homo sapiens

<400> 61
 Met Phe Leu Pro Thr Phe Tyr Pro Glu Lys Arg Phe Ser Pro Lys Asp
 1 5 10 15
 Ser Ala Gln Ser Val Pro Pro Trp Glu His Leu Pro Gly Gln Pro Leu
 20 25 30
 Arg Ala His Trp Ala Ser Leu His His Thr Asn Thr Pro Val Pro His
 35 40 45
 Trp Leu Ser Asp Tyr Met Ala Val Cys Leu Val Lys Lys Lys Asn Gln
 50 55 60
 Lys Lys Lys Lys Gln Lys Lys Lys Lys Lys Lys
 65 70 75

<210> 62
 <211> 93
 <212> PRT
 <213> Homo sapiens

<400> 62
 Val Gly Thr Ala Ile Met Glu Asn Ser Met Ala Val Pro Leu Lys Thr
 1 5 10 15
 Glu Leu Pro Tyr Asp Pro Ala Ile Pro Leu Leu Ser Ile Pro Lys Glu
 20 25 30
 Met Lys Ser Ala Leu His Arg Asp Ile Cys Ile Leu Met Leu Thr Ala
 35 40 45
 Ala Leu Phe Thr Ile Ala Lys Thr Glu Lys Gln His Lys Cys Pro Ser
 50 55 60

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<400> 64
Thr Arg Pro Val Leu Ala Tyr Val Leu Gly Asp Pro Ala Ile Tyr Gln
  1                      5                10              15
Ser Leu Lys Ala Gln Asn Ala Tyr Ser Arg His Cys Pro Phe Tyr Val
      20                25              30
Ser Ile Gln Ser Tyr Trp Leu Ser Phe Phe Met Val Met Ile Leu Phe

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45

Ser Ser Asp Asn Ala Gln Asn Phe Ser Phe Lys Thr Asn Ser Gly Phe

100 105 110
 Ala Ala Ala Ser Ser Gly Ser Pro Ala Gly Phe Gly Ser Ser Pro Ala
 115 120 125
 Phe Gly Ala Ala Ala Ser Thr Ser Ser Gly Ile Ser Thr Ser Ala Pro
 130 135 140
 Ala Phe Gly Phe Gly Lys Pro Glu Val Thr Ser Ala Ala Ser Phe Ser
 145 150 155 160
 Phe Lys Ser Pro Ala Ala Ser Ser Phe Gly Ser Pro Gly Phe Ser Gly
 165 170 175
 Leu Pro Ala Ser Leu Ala Thr Gly Pro Val Arg Ala Pro Val Ala Pro
 180 185 190
 Ala Phe Gly Gly Gly Ser Ser Val Ala Gly Phe Gly Ser Pro Gly Ser
 195 200 205
 His Ser His Thr Ala Phe Ser Lys Pro Ser Ser Asp Thr Phe Gly Asn
 210 215 220
 Ser Ser Ile Ser Thr Ser Leu Ser Ala Ser Ser Ser Ile Ile Ala Thr
 225 230 235 240
 Asp Asn Val Leu Phe Thr Pro Arg Asn Lys Leu Thr Val Glu Glu Leu
 245 250 255
 Glu Gln Phe Gln Ser Lys Lys Phe Thr Leu Gly Lys Ile
 260 265

 <210> 66
 <211> 300
 <212> PRT
 <213> Homo sapiens

 <400> 66
 Met Ser Ser Ser His Pro Val Ser Pro Asn Pro His His Gly Gly Ala
 1 5 10 15
 Ala Glu Ile Lys Lys Pro Asn Ile Ser Gly Phe Thr Asp Ile Ser Pro
 20 25 30
 Glu Glu Leu Arg Leu Glu Tyr His Asn Phe Leu Thr Ser Asn Asn Leu
 35 40 45
 Gln Ser Tyr Leu Asn Ser Val Gln Arg Leu Ile Asn Gln Trp Arg Asn
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Pro Ser Gly Asn Asn Arg Arg Gly Trp Asn Thr Thr Ser Gln Arg Tyr
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Ser Asn Val Ile Gln Pro Ser Ser Phe Ser Lys Ser Thr Pro Trp Gly
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TO THE "6212360"

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see p.5

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TETRA-6-ET-360

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8 <140> CURRENT APPLICATION NUMBER: US/09/832,129
9 <141> CURRENT FILING DATE: 2001-04-11
11 <150> PRIOR APPLICATION NUMBER: PCT/US00/28664
12 <151> PRIOR FILING DATE: 2000-10-17
14 <150> PRIOR APPLICATION NUMBER: 60/163,085
15 <151> PRIOR FILING DATE: 1999-11-02
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36 ggctgaatgg caaggagtac aagtgaagg tctccaacaa agccctccca acccccatcg      360
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38 catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggcttct      480
39 atccaagcga catgcgcgtg gagtgggaga gcaatgggca gccggagAAC aactacaaga      540
40 ccacgcctcc cgtgctggac tccgacggct ccttcttctt ctacagcaag ctcaccgtgg      600
41 acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggtctctg      660
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87	gcccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat	180
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 150 ctgctgacct gggagaatgg acgggttctg ggagaccaga tggctcaga cactgagctc 180
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288	cagttttgct	tgcattcctt	gagaatgtat	ttatctgaag	atcaaaaacaa	acaatccaga	3000
289	tgtataagta	ctaggcagaa	gccaatttta	aaatttcctt	gaataatcca	tgaaaggaat	3060
290	aattcaataa	cagataaaca	gagttggcag	tatattatag	tgataatttt	gtattttcac	3120
291	aaaaaaaaag	ttaaactcct	cttttctttt	tattataatg	accagctttt	ggtatttcat	3180

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/832,129

DATE: 04/27/2001

TIME: 13:10:54

Input Set : A:\PZ045Pl_SeqList04112001.txt

Output Set: N:\CRF3\04272001\I832129.raw

L:8 M:270 C: Current Application Number differs, Replaced Current Application Number
L:57 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:845 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:2556 M:283 W: Missing Blank Line separator, <400> field identifier
L:2566 M:283 W: Missing Blank Line separator, <400> field identifier

Total: 6272001